

<110> GENOMINE INC.  
KOREA RESEARCH INSTITUTE OF CHEMICAL TECHNOLOGY

<120> Polypeptide Participating in Pyridoxine Biosynthesis, a  
Polynucleotide Coding the Polypeptide and Those Uses

<150> KR 10-2004-0011517  
<151> 2004-02-20

<160> 6

<170> KopatentIn 1.71

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| aaatctctct tccattttct ccacacaaat ttctcttcaa tctccgataa tggaaggaac  | 120 |
| cggcgttggt gcggtgtacg gtaacgggtc gataacggag gogaagaaat ctcccttctc  | 180 |
| cgtgaaggtc ggtttggctc agatgctccg tgggtggttt atcatggatg tcgtcaacgc  | 240 |
| cgagcaagct cgtatcgccg aggaggctgg tgcttgcgcc gtcattggctt tggagcgtgt | 300 |
| tcctgctgat atccgcgctc aaggaggcgt cgctcgtatg agcgatccac aaatgattaa  | 360 |

|  |      |
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| agaaatcaaa caagccgtta cgattccggt gatggctaag gctaggattg gtcatttcgt  | 420  |
| tgaagctcag atccttgaag caattggaat cgattacatc gatgagagcg aggttttgac  | 480  |
| tcttgctgat gaagatcatc acatcaacaa gcataatttc cggatcccg tctgttcggg   | 540  |
| ttgccggaat ctcggcgagg ctctgaggag gatccgtgaa ggtcgggcga tgattaggac  | 600  |
| caaaggtgaa gctggaaccg gtaacattat tgaagctgtg aggcattgtga ggtctgttaa | 660  |
| tggtagacatt agggttttgc gaaacatgga tgatgatgag gttttcactt tcgctaagaa | 720  |
| attagccgct ccgtacgac tcgtgatgca gactaagcag ctgggtcgtc ttctgtagt    | 780  |
| ccaattcgcc gccggtggag tggctactcc ggctgatgca gctctcatga tgcagcttgg  | 840  |
| atgtgatggt gtctttgttg gttctggtat cttcaagagc ggtgaccag ctgctcgtgc   | 900  |
| acgtgccatt gttcaggctg tgactcatta cagtgacct gagatgcttg tggaggtag    | 960  |
| ctgtgggcct ggagaagcca tggttgggat caatctcaac gatgagaagg ttgagaggtt  | 1020 |
| cgctaatcgc tccgagtgat caaagaaata aaaggtaaaa tatctcagac gaaatggttt  | 1080 |
| cagaattttc tcagaccatt ttgcagtaat ctctttgaaa agaagaagat gatgatattg  | 1140 |
| ttggtagttt gtatcctttg tgttttcctt ataatctttg atagtctttt gttattgtaa  | 1200 |
| ctcgtaatcc ctttgcaaga acaagtttgt cagttataat aatgtactac tctcttgatc  | 1260 |

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Leu Arg Gly Gly Val Ile Met Asp Val Val Asn Ala Glu Gln Ala Arg  
35 40 45

Ile Ala Glu Glu Ala Gly Ala Cys Ala Val Met Ala Leu Glu Arg Val  
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Pro Ala Asp Ile Arg Ala Gln Gly Gly Val Ala Arg Met Ser Asp Pro  
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Gln Met Ile Lys Glu Ile Lys Gln Ala Val Thr Ile Pro Val Met Ala  
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Lys Ala Arg Ile Gly His Phe Val Glu Ala Gln Ile Leu Glu Ala Ile  
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Gly Ile Asp Tyr Ile Asp Glu Ser Glu Val Leu Thr Leu Ala Asp Glu  
115 120 125

Asp His His Ile Asn Lys His Asn Phe Arg Ile Pro Phe Val Cys Gly  
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Cys Arg Asn Leu Gly Glu Ala Leu Arg Arg Ile Arg Glu Gly Ala Ala  
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Met Ile Arg Thr Lys Gly Glu Ala Gly Thr Gly Asn Ile Ile Glu Ala  
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Val Arg His Val Arg Ser Val Asn Gly Asp Ile Arg Val Leu Arg Asn  
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Met Asp Asp Asp Glu Val Phe Thr Phe Ala Lys Lys Leu Ala Ala Pro  
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Tyr Asp Leu Val Met Gln Thr Lys Gln Leu Gly Arg Leu Pro Val Val  
210 215 220

Gln Phe Ala Ala Gly Gly Val Ala Thr Pro Ala Asp Ala Ala Leu Met  
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Met Gln Leu Gly Cys Asp Gly Val Phe Val Gly Ser Gly Ile Phe Lys  
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Ser Gly Asp Pro Ala Arg Arg Ala Arg Ala Ile Val Gln Ala Val Thr  
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His Tyr Ser Asp Pro Glu Met Leu Val Glu Val Ser Cys Gly Leu Gly  
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